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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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    US-09-027-337-1
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US-09-644-600-1
US-09-635-008-39
US-09-535-008-39
US-09-535-008-39
US-09-644-600-9
US-09-252-991A-8158
US-08-650-766-2
US-09-389-487-2
US-08-922-635-2
US-08-922-635-3
US-08-922-635-3
US-08-922-635-1
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US-09-252-991A-14671
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US-08-943-731-640
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Patent No. 5972616

GRNERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
APPLICANT: Tanimoto, Hirotoshi
TITLE OF INVENTION: TADO-15: An Extracellular Serine Protease Overexpressed in TITLE OF INVENTION: Breast and Ovarian Carcinomas
FILE REFERENCE: D6046

CURRENT APPLICATION NUMBER: US/09/027,337B

CURRENT FILING DATE: 1998-02-20

NUMBER OF SEQ ID NOS: 13

SEQ ID NO 1

LENGTH: 3147
TYPE: DNA

ORGANISM: Homo sapiens
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; LOCATION: 23..2589
; OTHER INFORMATION: cDNA sequence of TADG-15
US-09-027-337-1
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US-09-027-337-1/c
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Best Local S
Matches 159
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                                                                                                  TATGTTCTATGATGTGCCTGAAGAACAGGACCTGTACAACAAGTTCAAGGGACGCGTGC
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US-08-99-126-280-3
US-09-126-280-1
US-09-126-280-1
US-09-277-858-1
US-08-340-011-1
US-08-340-011-1
US-08-340-011-1
US-08-795-430-1
US-08-795-700-1
US-08-795-700-1
US-08-795-700-1
US-08-795-701-3
US-08-946-648-31
US-08-446-648-31
US-08-340-011-3
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US-08-340-011-3
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Pred. No. 0.041
0; Mismatches
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Scoring table: Sequence: Title: Perfect score:

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Sequence 18, Application US/09644600

Patent No. 6451500

GENERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
APPLICANT: Tanimoto, Hirotoshi
ITILE OF INVENTION: TADG-15: An Extracellular Serine Protease
ITILE OF INVENTION: O'Overexpressed in Carcinomas
FILE REFERENCE: D6064CIP/D

CURRENT FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: US/09/644,600

CURRENT FILING DATE: 1999-10-20
PRIOR PILING DATE: 1999-10-20
PRIOR PILING DATE: 1999-10-20
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SEQ ID NO 1
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Best Local Similarity 47.9%;
Matches 159; Conservative
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Patent No. 6451500
GENERAL INFORMATION:
                                 PRIOR APPLICATION NUMBER: 09/027,337 PRIOR FILING DATE: 1998-02-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/644,600
CURRENT FILING DATE: 2000-08-23
PRIOR APPLICATION UNMBER: 09/421,213
PRIOR FILING DATE: 1999-10-20
PRIOR PILING DATE: 1998-02-20
PRIOR FILING DATE: 1998-02-20
INTERIOR DATE: 1998-02-20
INTERIOR SEQ ID NOS: 98
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TITLE OF INVENTION: TADG-15: An Extracellular Serine Protease
TITLE OF INVENTION: Overexpressed in Carcinomas
FILE REPERENCE: D6064CIP/D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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                 SEQ ID NOS:
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; TYPE: RNA
; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: Antisense of TADG-15
US-09-644-600-18
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US-08-232-463-14/c
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Best Local Similarity 39.8%;
Matches 132; Conservative 2'
                                                                                                   APPLICATION NUMBER: US/07/
FILLING DATE:
FILING DATE: EP 91
APPLICATION NUMBER: EP 91
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
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CITY: Alexandria
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APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
APPLICANT: FALKNER, F. G.
APPLICANT: FALKNER, F. G.
APPLICANT: FALKNER, F. G.
APPLICANT: PALKNER, F. G.
APPLICANT: SCHEME PALKNER, F. G.
APPLICANT: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30.
'ELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            esaceerricicceaeciccaaercccccice ззв
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGGCCCCGTCTGGCCCCCCAGAATGTGACGCTGCTCTCCCAGAACTTCAGCGTGTAC 126
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1800 Diagonal Road,
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                                                                                                                                                                                                                                                             EP 91 114 300.6
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                                                                      30472/114 IMMU
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Pred. No. 0.041;
7; Mismatches 171;
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TELEFAX: (703)683-4109
TELEX: 899149
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:

14:

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; NUMBER OF SEQ ID NOS: 77
; SOFTWARE: Patentin Ver. 2
; SEQ ID NO 39
; LENGTH: 971
; TYPE: DNA
; ORGANISM: Homo Bapiens
US-09-535-008-39
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                                Query Match 2.6
Best Local Similarity 49.3
Matches 110; Conservative
                                                                                                                                                                                                                   FILE REFERENCE: 2318-259
CURRENT APPLICATION NUMBER: US/09/535,008
CURRENT FILING DATE: 2000-03-23
EARLIER APPLICATION NUMBER: U.S. 60/125,806
EARLIER FILING DATE: 1999-03-23
NUMBER OF SEQ ID NOS: 77
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                                                                                                                                                                                                                                                                                                               APPLICANT: Wong, Alexander K.C.
APPLICANT: Tavtigian, Sean V.
APPLICANT: Tavtigian, Sean V.
APPLICANT: Tool, David H.-F.
TITLE OF INVENTION: BRGI IS A TUMOR SUPPRESSOR THAT IS
TITLE OF INVENTION: AND OTHER CANCER TYPES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .09-535-008-39/c
                                                                                                                                                                                                                                                                                                                                                                                                                                  equence 39, Application US/09535008 atent No. 6465629 ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             / Match 2.7%; Score 42.6; DB 1; Local Similarity 3.8%; Pred. No. 0.17; DB 1; 15; Conservative 212; Mismatches 166;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 7218 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
1171 GCTGGGCCAGCACTGTGGACTCCTCCTGGGACAGGGCTGGGTCCTCTGGCTATTTGGCTG 1230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1200 GACAGGGCTGGGTCCTCTGGCTATTTGGCTGAGAAGGGCCCAGGCCAAGGGCCGGGTGGG 1255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1080 CACTCGGAGGCTGGTGGGGTGGACTCAGGGAGGCCCAGGGCTCCTCTGGTCCCAAGCGAA 1139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1076 RRRRRRRRRRATCGCAAGCTCCCTCGACCTGCA 1044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1260 GATGGGCACCAAGAATCTCTCCCACCACCTGAA 1292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       960 AAGAAGGACCTTGCAGAGGACGAAGAGGAGGAGGATGAGGAGGACACAGAAGATGGCGTC 1019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    900 ACCAGAGGGGTCAGGCCGACGCCTCGAGTCAGGGCCCAGCCCAACAGACAAGATGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGCTCCTCTGCTTGGGATTCTTCAGACAGAAGCTGGGCCAGCACTGTGGACTCCTCCTGG 1199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGCTTCCAGCCCTACATTGAACCACCTTCTTTCCTGGGGCAAGAGCACCAGGCTCCAGGG 1079
                                                     2.6%;
49.3%;
                                Score 42.2; DB 4;
Pred. No. 0.1;
0; Mismatches 113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 7218;
                                                                       Length 971;
                                    Indels
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US-09-027-337-9/c
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Sequence 9, Application US/09644600
Patent No. 6431500
GENERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
APPLICANT: Tanimoto, Hirotoshi
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Best Local Similarity 48.3%;
Matches 117; Conservative
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Patent No. 5972616
GENERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
APPLICANT: Tanimoto, Hirotoshi
TITLE OF INVENTION: TADG-15: An Extracellular Serine Protease Overexpressed in
TITLE OF INVENTION: Breast and Ovarian Carcinomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 9
LENGTH: 2900
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CURRENT FILING DATE: 1998-02-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              323
                                                                                                                                                                                                                                                                          203
                                                                                                                                                                                                                                                                                                                                                                                                127 CTGACATGGCTCCCAGGGCTTGGCAACCCCCAGGATGTGACCTATTTTGTGGCCCATCAGA 186
                                                                                                                                                                                                                                                                                                                                                                                                                                         263 GACTCCTTGTGGTAGGGGCCCAGGAATGGGACTCCGCTGTACAGCAGCTTCAGCGCGTCC
                                                                                                                                                                                                                                 247 TA 248
                                                                                                                                                                                                                                                                                                             187 GCTCTCCCACCCGTAGACGGTGGCGCGAAGTGGAAGAGTGTGCGGGAACCAAGGAGCTGC 246
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                                                                                                                                                                                        83 TA 82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7 ATGGCGGGGCCCGAGGGCTGCCCCTGCTCCTGTGCCTGCAGGCCGCTCCAGGG 66
                                                                                                                                                                                                                                                                                                                                                         TTCACCTTGCTGGCCAGGCTTACAAACTCAGTGGAGTTGGAGTTCTCGTAGGCATCCACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATGCTGAACTCAGACCAGTAGTAGGCGATGACGCTGCCCTCGCTGAAGGCCCGTCACAGCC 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AATTCTCCAAGGACTCGGGTTTCCTGGAAGAGCTCCCAGAAGATAACCTCTCCTCCTGGG 1350
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Pred. No. 0.17;
0; Mismatches 125; Indels
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APPLICANT: Tanimoto, Hirotoshi TITLE OF INVENTION: TADG-15: An Extracellular Serine Protease TITLE OF INVENTION: Overexpressed in Carcinomas FILE REFERENCE: D6064CIP/D

CURRENT APPLICATION NUMBER: US/09/644,600

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RESULT 8
US-09-252-991A-8224
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                                                                                                                                      FEATURE:
NAME/KEY: unsure
LOCATION: (948)
JOTHER IMPORMATION: Identity of nucleotide at the above locations are unknown.
US-09-252-991A-8224
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; ORGANISM: Homo sapiens
; PEATURE:
; OTHER INFORMATION: SNC-19; GeneBank Accession No.
US-09-644-600-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 09/421,213
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: 09/027,337
PRIOR FILING DATE: 1998-02-20
NUMBER OF SEQ ID NOS: 98
SEQ ID NO 9
LENGTH: 2900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 6551795
GENERAL INFORMATION:
                                                                                  Query Match 2.6%;
Best Local Similarity 46.5%;
Matches 134; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 8224, Application US/09252991A
Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                APPLICANT: MATC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US/60/074,788
PRIOR PILING DATE: 1998-02-18
PRIOR FILING DATE: US/60/094,190
RIOR APPLICATION NUMBER: US/60/094,190
RIOR APPLICATION NUMBER: US/60/094,190
RIOR APPLICATION NUMBER: US/60/094,190
RIOR APPLICATION NUMBER: US/60/094,190
RIOR APPLICATION DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
LENGTH: 975
TYPE: NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches 117;
                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Pseudomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             247
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                                                 843 CAGCCCAGCAGACCAGAGTCCGTGAATGACTTGTTCCTCTGTCCCCAAAAAGGAACTGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               263 GACTCCTTGTGGTAGGGGCCCAGGAATGGGACTCCGCTGTACAGCAGCTTCAGCGCGTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  67
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                  CGGCCGCGATCACCACAGGCCGCCATGCACGCATTCCCCCCTCACCCTGGACAACCGACTC
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illarity 48.3%;
Conservative
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                                                                               Score 41.6; DB 4;
Pred. No. 0.14;
0; Mismatches 154;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 42; DB'4; Length 2900 Pred. No. 0.17;
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                                                                                                                Length 975;
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US-08-650-766-2
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Sequence 2, Application US/08650766D Patent No. 6015690 GENERAL INFORMATION:
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RESULT 9
US-09-252-991A-8158
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; NAME/KEY: unsure
; LOCATION: (982)
; LOCATION: 1982)
; OTHER INFORMATION: Identity of nucleotide at the above locations are unknown.
US-09-252-991A-8158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8158, Applic Patent No. 6551795 GENERAL INFORMATION:
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Best Local Similarity
Matches 134; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: MATC J. Rubenfield et al.
TITLE OF INVENTION: NUCLBIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196, 136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILLING DATE: 1999-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION UNMER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION UNMER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Pseudomonas aeruginosa
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                                                                                                                                1023
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                                                                                                                   TTCCAGCCCTACATTGAACCACCTTCTTTCCTGGGGCAAGAGCACCAGGCTCCCAGGGCAC 1082
                                        ATCGACCTCGCCGACTGGAAAGCCGACTGGCGCGAACTGATCGCCGCGCGCTCGACAGGTTT 280
                                                                                                                                                                                                                                                                                                                                                               CGGCCGCGATCACCACAGGCCGCCATGCACGCATTCCCCCCTCACCCTGGACAACCGACTC
                                                                                                                                                                                                                                                                                                                                                                                                          CAGCCCAGCAGACCAGAGTCCGTGAATGACTTGTTCCTCTGTCCCCAAAAGGAACTGACC
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GCCTGGGCCTGGGGCCAGGCCGCGAAAAACGCCAGTACGCTGCTGCTC 388
                                                                                                                                                                                                                                                                     TCGGAGGCTGGTGGACTCAGGGAGGCCCAGGGCTCCTCTGGTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         154;
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Best Local Similarity
"""has 75; Conserve
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; ORGANISM: Homo sapiens
US-09-389-487-2
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; ORGANISM: Homo sapiens
US-08-650-766-2
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                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 6576742
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NUMBER OF SEQ ID NOS: 21
SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/389,487
CURRENT FILING DATE: 1999-09-03
EARLIER APPLICATION NUMBER: US 08/650,766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: PILETZ, John B.
APPLICANT: IVANOV, Tina R.
APPLICANT: IVANOV, Tina R.
TITLE OF INVENTION: DAM SEQUENCE ENCODING A HUMAN IMIDAZOLINE RECEPTOR AND
TITLE OF INVENTION: METHOD FOR CLONING THE SAME
FILE REFERENCE: Corrected Sequence Listing
                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   atent No. 601569
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APPLICANT: IVANOV, TIMA R.
TITLE OF INVENTION: DNA SEQUENCE ENCODING A HUMAN IMIDAZOLINE RECEPTOR AND
TITLE OF INVENTION: METHOD FOR CLONING THE SAME
FILE REFERENCE: Corrected Sequence Listing
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tent No. 6576742
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                               TGGTGGGGTGGA 1102
                                                                CTACTTTGAAATGGGGCCCCCAGACGTGGAGGAGGAGGAGGAGGAGGCCAGGGGGAGGA
                                                                                                 CTACATTGAACCACCTTCTTTCCTGGGGCAAGAGCACCAGGCTCCAGGGCACTCGGAGGC 1090
                                                                                                                                                                       TGCAGAGGACGAAGAGGAGGAGGATGAGGAGGAGACACAGAAGATGGCGTCAGCTTCCAGCC 1030
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AGAGGAGGAGGA 794
                                                                                                                                    TGAACAGGGCGAGGAGGATGAGGAGGAGGAAGAAGAGGAGGACGTGGCTGAGAACCG 722
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ilarity 56.8%;
Conservative
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                                                                                                                                                                                                                       Score 40.8; DB 4;
Pred. No. 0.29;
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                                                Matches
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Best Local &
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RESULT 12
US-08-922-635-2
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                                                                                                                                                                          NUMBER OF SEQ ID NOS: 2:
SOFTWARE: PatentIn Ver.
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                            APPLICANT: PILETZ, John B.

APPLICANT: IVANOY, Tina R.

TITLE OF INVENTION: DNA SEQUENCE ENCODING A HUMAN IMIDAZOLINE RECEPTOR TITLE OF INVENTION: METHOD FOR CLONING THE SAME FILE REFERENCE: COFFECTE SEQUENCE LISTING SAME

FILE REFERENCE: COFFECTE SEQUENCE LISTING

Patent No. 6015690
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3, Application US/08650766D Patent No. 6015690 GENERAL INFORMATION:
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SENERAL INCORNATION:

APPLICANT: PILETZ, John B.

APPLICANT: ITANOV, Tina R.

TITLE OF INVENTION: DAM MOLECULES ENCODED THEREBY

TITLE OF INVENTION: AND POLYPEPTIDES ENCODED THEREBY

FILE REFERENCE: Corrected Sequence Listing
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LENGTH: 1954
TYPE: DNA
ORGANISM: Homo sapiens
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Patent No. 60338
                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
-08-650-766-3
                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/08/650,766D CURRENT FILING DATE: 1996-05-20 EARLIER APPLICATION NUMBER: US 60/012,600 EARLIER FILING DATE: 1996-03-01 NUMBER OF SEQ ID NOS: 21
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CURRENT FILING DATE: 1997-09-03
EARLIER APPLICATION NUMBER: 08/650,766
                                                                                                                                      LENGTH:
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  Score 40.8; DB 3; Pred. No. 0.38; 0; Mismatches 57;
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SOFTWARE: PATENTIN Ver. 2
SEQ ID NO 3
LENGTH: 3318
TYPE: DNA
ORGANIZM: Homo sapiens
US-08-922-635-3
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US-08-922-635-3
                               CURRENT APPLICATION NUMBER: US/09/389,487
CURRENT FILING DATE: 1999-09-03
EARLIER APPLICATION NUMBER: US 08/650,766
EARLIER FILING DATE: 1996-05-20
NUMBER OF SEQ ID NOS: 21
SOFTWARE: PACENTIN Ver. 2.0
SEQ ID NO 3
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Best Local Similarity 56.8%;
Matches 75; Conservative
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APPLICANT: IVANOV, TINB R.
TITLE OF INVENTION: DNA SEQUENCE ENCODING A HUMAN
TITLE OF INVENTION: METHOD FOR CLONING THE SAME
PILE REFERENCE: Corrected Sequence Listing
Patent No. 6576742
                                                                                                                                                                                                                                                                                                                             Sequence 3, Application US/09389487 Patent No. 6576742
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APPLICANT: IVANDOV, TIAR R.
TITLE OF INVENTION: DNA MOLECULES ENCODING IMIDALINE RECEPTIVE POLYPEPTIDES
TITLE OF INVENTION: AND POLYPEPTIDES ENCODED THEREBY
TYPE: DNA
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US-09-389-487-3
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Best Local Similarity 56.8%;
Matches 75; Conservative
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